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RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/050,902

DATE: 01/23/2003

TIME: 12:20:04

Input Set : A:\1700.0190004seqlist.txt
Output Set: N:\CRF4\01232003\J050902.raw

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5 <110> APPLICANT: Renner, Wolfgang A.
         Bachmann, Martin
 6
         Tissot, Alain
 7
         Maurer, Patrick
 8
         Lechner, Franziska
9
10
         Sebbel, Peter
         Piossek, Christine
11
13 <120> TITLE OF INVENTION: Molecular Antigen Array
15 <130> FILE REFERENCE: 1700.0190004
17 <140> CURRENT APPLICATION NUMBER: 10/050,902
18 <141> CURRENT FILING DATE: 2002-01-18
20 <150> PRIOR APPLICATION NUMBER: US 60/262,379
21 <151> PRIOR FILING DATE: 2001-01-19
23 <150> PRIOR APPLICATION NUMBER: US 60/288,549
24 <151> PRIOR FILING DATE: 2001-05-04
26 <150> PRIOR APPLICATION NUMBER: US 60/326,998
27 <151> PRIOR FILING DATE: 2001-10-05
29 <150> PRIOR APPLICATION NUMBER: US 60/331,045
30 <151> PRIOR FILING DATE: 2001-11-07
32 <160> NUMBER OF SEQ ID NOS: 350
34 <170> SOFTWARE: PatentIn Ver. 2.1
36 <210> SEQ ID NO: 1
37 <211> LENGTH: 41
38 <212> TYPE: DNA
39 <213> ORGANISM: Artificial Sequence
41 <220> FEATURE:
42 <223> OTHER INFORMATION: Primer
44 <400> SEQUENCE: 1
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48 <210> SEQ ID NO: 2
49 <211> LENGTH: 44
50 <212> TYPE: DNA
51 <213> ORGANISM: Artificial Sequence
53 <220> FEATURE:
54 <223> OTHER INFORMATION: Primer
56 <400> SEQUENCE: 2
57 cggtggttac ctgctgcacg cgttgcttaa gcgacatgta gcgg
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61 <211> LENGTH: 20
62 <212> TYPE: DNA
63 <213> ORGANISM: Artificial Sequence
65 <220> FEATURE:
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66 <223> OTHER INFORMATION: Primer

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68 <400> SEQUENCE: 3 20 69 ccatgaggcc tacgataccc 72 <210> SEQ ID NO: 4 73 <211> LENGTH: 25 74 <212> TYPE: DNA 75 <213> ORGANISM: Artificial Sequence 77 <220> FEATURE: 78 <223> OTHER INFORMATION: Primer 80 <400> SEQUENCE: 4 25 81 ggcactcacg gcgcgcttta caggc 84 <210> SEQ ID NO: 5 85 <211> LENGTH: 47 86 <212> TYPE: DNA 87 <213> ORGANISM: Artificial Sequence 89 <220> FEATURE: 90 <223> OTHER INFORMATION: Primer 92 <400> SEQUENCE: 5 47 93 ccttctttaa cggtggttac ctgctggcaa ccaacgtggt tcatgac 96 <210> SEQ ID NO: 6 97 <211> LENGTH: 40 98 <212> TYPE: DNA 99 <213> ORGANISM: Artificial Sequence 101 <220> FEATURE: 102 <223> OTHER INFORMATION: Primer 104 <400> SEQUENCE: 6 40 105 aagcatgctg cacgcgtgtg cggtggtcgg atcgcccggc 108 <210> SEQ ID NO: 7 109 <211> LENGTH: 90 110 <212> TYPE: DNA 111 <213> ORGANISM: Artificial Sequence 113 <220> FEATURE: 114 <223> OTHER INFORMATION: Primer 116 <400> SEQUENCE: 7 117 gggtctagat tcccaaccat tcccttatcc aggctttttg acaacgctat gctccgcgcc 60 118 catcgtctgc accagctggc ctttgacacc 121 <210> SEQ ID NO: 8 122 <211> LENGTH: 108 123 <212> TYPE: DNA 124 <213> ORGANISM: Artificial Sequence 126 <220> FEATURE: 127 <223> OTHER INFORMATION: Primer 129 <400> SEQUENCE: 8 130 gggtctagaa ggaggtaaaa aacgatgaaa aagacagcta tcgcgattgc agtggcactg 60 131 gctggtttcg ctaccgtagc gcaggccttc ccaaccattc ccttatcc 108 134 <210> SEQ ID NO: 9 135 <211> LENGTH: 31 136 <212> TYPE: DNA 137 <213> ORGANISM: Artificial Sequence 139 <220> FEATURE:

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140 <223> OTHER INFORMATION: Primer
142 <400> SEQUENCE: 9
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143 cccgaattcc tagaagccac agctgccctc c
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147 <211> LENGTH: 24
148 <212> TYPE: DNA
149 <213> ORGANISM: Artificial Sequence
151 <220> FEATURE:
152 <223> OTHER INFORMATION: Primer
154 <400> SEQUENCE: 10
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158 <210> SEQ ID NO: 11
159 <211> LENGTH: 41
160 <212> TYPE: DNA
161 <213> ORGANISM: Artificial Sequence
163 <220> FEATURE:
164 <223> OTHER INFORMATION: Primer
166 <400> SEQUENCE: 11
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167 ccgcggaaga gccaccgcaa ccaccgtgtg ccgccaggat g
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171 <211> LENGTH: 33
172 <212> TYPE: DNA
173 <213> ORGANISM: Artificial Sequence
175 <220> FEATURE:
176 <223> OTHER INFORMATION: Primer
178 <400> SEQUENCE: 12
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179 ctatcatcta gaatgaatag aggattcttt aac
182 <210> SEQ ID NO: 13
183 <211> LENGTH: 15
184 <212> TYPE: DNA
185 <213> ORGANISM: Artificial Sequence
187 <220> FEATURE:
188 <223> OTHER INFORMATION: Modified ribosome binding site
190 <400> SEQUENCE: 13
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191 aggaggtaaa aaacg
194 <210> SEQ ID NO: 14
195 <211> LENGTH: 21
 196 <212> TYPE: PRT
 197 <213> ORGANISM: Artificial Sequence
 199 <220> FEATURE:
 200 <223> OTHER INFORMATION: signal peptide
 202 <400> SEQUENCE: 14
203 Met Lys Lys Thr Ala Ile Ala Ile Ala Val Ala Leu Ala Gly Phe Ala
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                       5
 204
     1
 206 Thr Val Ala Gln Ala
 207
 210 <210> SEQ ID NO: 15
 211 <211> LENGTH: 46
 212 <212> TYPE: PRT
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213 <213> ORGANISM: Artificial Sequence
215 <220> FEATURE:
216 <223> OTHER INFORMATION: modified Fos construct
218 <400> SEQUENCE: 15
219 Cys Gly Gly Leu Thr Asp Thr Leu Gln Ala Glu Thr Asp Gln Val Glu
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220
     1
222 Asp Glu Lys Ser Ala Leu Gln Thr Glu Ile Ala Asn Leu Leu Lys Glu
                                     25
                 20
225 Lys Glu Lys Leu Glu Phe Ile Leu Ala Ala His Gly Gly Cys
                                 40
226
             35
229 <210> SEQ ID NO: 16
230 <211> LENGTH: 6
231 <212> TYPE: PRT
232 <213> ORGANISM: Artificial Sequence
234 <220> FEATURE:
235 <223> OTHER INFORMATION: peptide linker
237 <400> SEQUENCE: 16
238 Ala Ala Ala Ser Gly Gly
242 <210> SEQ ID NO: 17
243 <211> LENGTH: 6
244 <212> TYPE: PRT
245 <213> ORGANISM: Artificial Sequence
247 <220> FEATURE:
248 <223> OTHER INFORMATION: peptide linker
250 <400> SEQUENCE: 17
251 Gly Gly Ser Ala Ala Ala
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255 <210> SEQ ID NO: 18
256 <211> LENGTH: 256
257 <212> TYPE: DNA
258 <213> ORGANISM: Artificial Sequence
260 <220> FEATURE:
261 <223> OTHER INFORMATION: Fos fusion construct
263 <400> SEQUENCE: 18
264 gaattcagga ggtaaaaaac gatgaaaaag acagctatcg cgattgcagt ggcactggct 60
265 ggtttcgcta ccgtagcgca ggcctgggtg ggggcggccg cttctggtgg ttgcggtggt 120
266 ctgaccgaca ccctgcaggc ggaaaccgac caggtggaag acgaaaaatc cgcgctgcaa 180
267 accgaaatcg cgaacctgct gaaagaaaaa gaaaagctgg agttcatcct ggcggcacac 240
                               256
268 ggtggttgct aagctt
271 <210> SEQ ID NO: 19
272 <211> LENGTH: 52
273 <212> TYPE: PRT
274 <213> ORGANISM: Artificial Sequence
276 <220> FEATURE:
 277 <223> OTHER INFORMATION: Fos fusion construct
 279 <400> SEQUENCE: 19
280 Ala Ala Ser Gly Gly Cys Gly Gly Leu Thr Asp Thr Leu Gln Ala
                       5
                                           10
 281
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283 Glu Thr Asp Gln Val Glu Asp Glu Lys Ser Ala Leu Gln Thr Glu Ile
                                     25
                 20
286 Ala Asn Leu Leu Lys Glu Lys Glu Lys Leu Glu Phe Ile Leu Ala Ala
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289 His Gly Gly Cys
         50
290
294 <210> SEQ ID NO: 20
295 <211> LENGTH: 261
296 <212> TYPE: DNA
297 <213> ORGANISM: Artificial Sequence
299 <220> FEATURE:
300 <223> OTHER INFORMATION: Fos fusion construct
303 <220> FEATURE:
304 <221> NAME/KEY: CDS
305 <222> LOCATION: (22)..(240)
307 <400> SEQUENCE: 20
308 gaattcagga ggtaaaaaac g atg aaa aag aca gct atc gcg att gca gtg
                            Met Lys Lys Thr Ala Ile Ala Ile Ala Val
309
310
312 gca ctg gct ggt ttc gct acc gta gcg cag gcc tgc ggt ggt ctg acc
                                                                        99
313 Ala Leu Ala Gly Phe Ala Thr Val Ala Gln Ala Cys Gly Gly Leu Thr
                                          20
316 gac acc ctg cag gcg gaa acc gac cag gtg gaa gac gaa aaa tcc gcg
                                                                        147
317 Asp Thr Leu Gln Ala Glu Thr Asp Gln Val Glu Asp Glu Lys Ser Ala
318
320 ctg caa acc gaa atc gcg aac ctg ctg aaa gaa aaa gaa aag ctg gag
                                                                        195
321 Leu Gln Thr Glu Ile Ala Asn Leu Leu Lys Glu Lys Glu Lys Leu Glu
                                  50
             45
324 ttc atc ctg gcg gca cac ggt ggt tgc ggt ggt tct gcg gcc gct
                                                                        240
325 Phe Ile Leu Ala Ala His Gly Gly Cys Gly Gly Ser Ala Ala Ala
                                                  70
                              65
         60
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328 gggtgtgggg atatcaagct t
331 <210> SEQ ID NO: 21
332 <211> LENGTH: 73
333 <212> TYPE: PRT
334 <213> ORGANISM: Artificial Sequence
336 <220> FEATURE:
337 <223> OTHER INFORMATION: Fos fusion construct
339 <400> SEQUENCE: 21
340 Met Lys Lys Thr Ala Ile Ala Ile Ala Val Ala Leu Ala Gly Phe Ala
343 Thr Val Ala Gln Ala Cys Gly Gly Leu Thr Asp Thr Leu Gln Ala Glu
                 20
346 Thr Asp Gln Val Glu Asp Glu Lys Ser Ala Leu Gln Thr Glu Ile Ala
                                  40
349 Asn Leu Leu Lys Glu Lys Glu Lys Leu Glu Phe Ile Leu Ala Ala His
352 Gly Gly Cys Gly Gly Ser Ala Ala Ala
353 65
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RAW SEQUENCE LISTING ERROR SUMMARY PATENT APPLICATION: US/10/050,902 DATE: 01/23/2003 TIME: 12:20:05

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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:111; Xaa Pos. 28 Seq#:283; N Pos. 9872

VERIFICATION SUMMARY

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DATE: 01/23/2003 TIME: 12:20:05

Input Set : A:\1700.0190004seqlist.txt
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L:2382 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:111 after pos.:16

L:5203 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (177) SEQUENCE:

L:9586 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:283 after pos.:9840